GenCore version 6.2.1 Copyright (c) 1993 - 2007 Biocceleration Ltd.

OM nucleic - nucleic search, using sw model

Run on:

June 14, 2007, 04:31:56; Search time 32216

Seconds

(without alignments) 10931.438 Million cell

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Total number of hits satisfying chosen parameters: 15137082

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Post-processing: Minimum Match 0%

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Listing first 45 summaries

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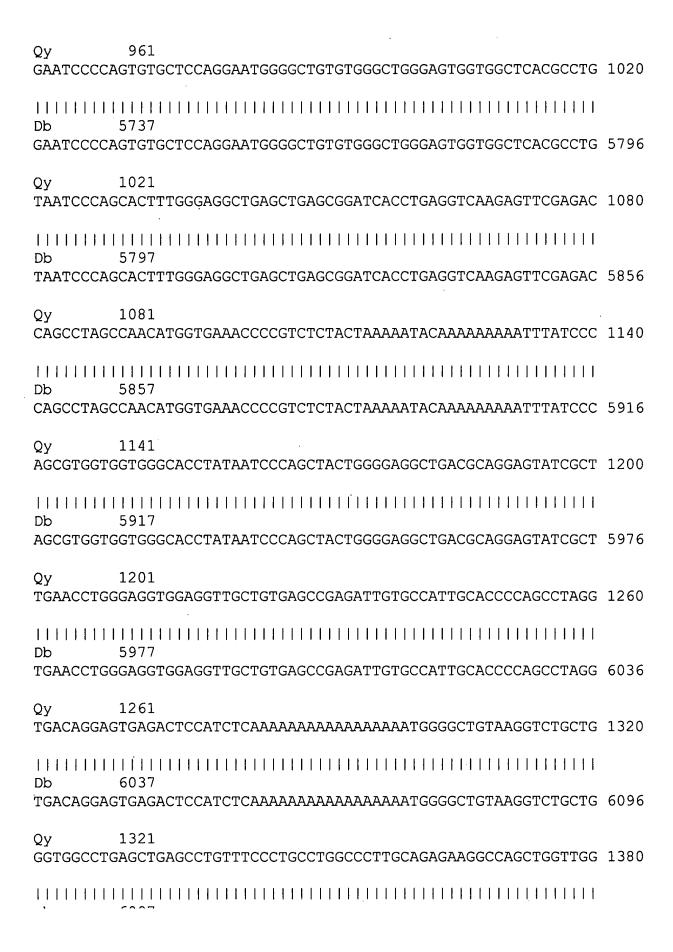
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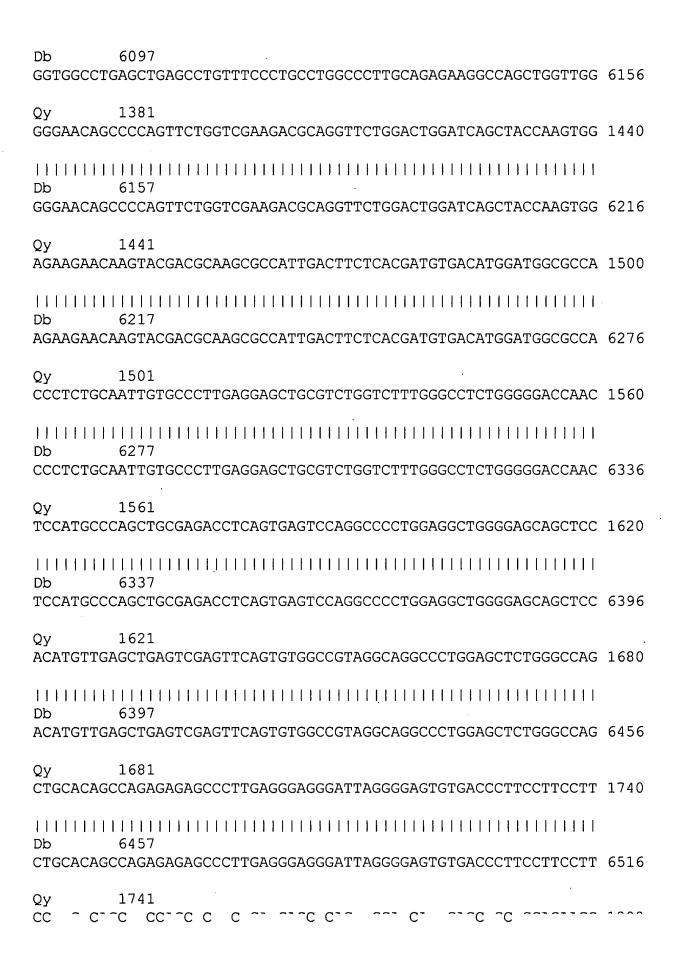
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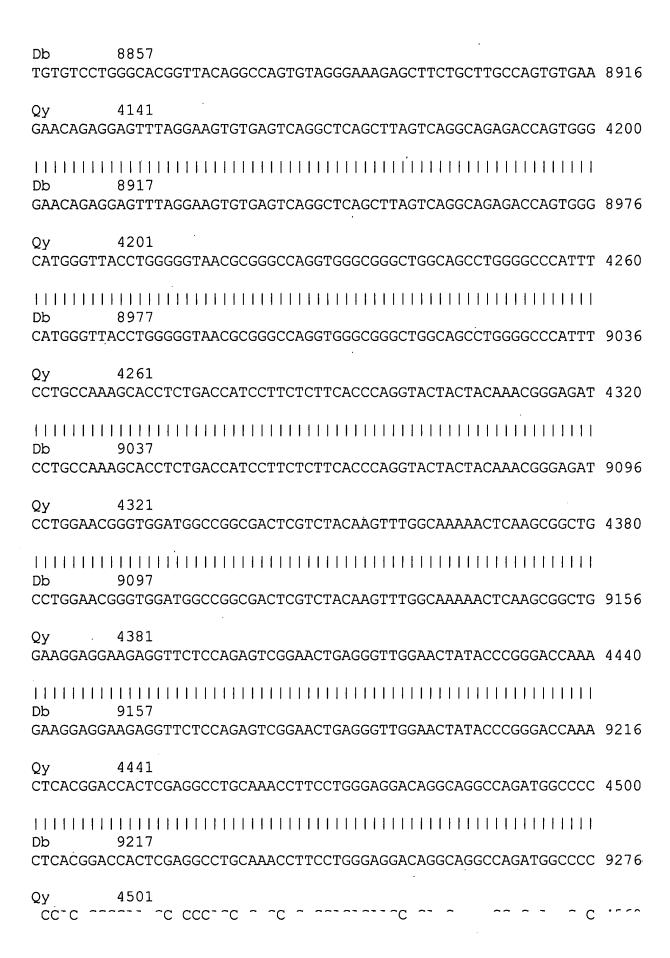
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GenCore version 6.2.1 Copyright (c) 1993 - 2007 Biocceleration Ltd.

OM nucleic - nucleic search, using sw model

Run on:

June 14, 2007, 04:31:56; Search time 854 Seconds

(without alignments) 10931.438 Million cell

updates/sec

Title:

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Scoring table:

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Total number of hits satisfying chosen parameters:

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Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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and Benz, C.C.
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Score 131; DB 5; Length 10772;

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LOCUS
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PRI 18-MAY-2005
DEFINITION Human DNA sequence from clone RP11-510N19 on
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transcription
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(CRIP1)
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KEYWORDS
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SOURCE
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;

GenCore version 6.2.1 Copyright (c) 1993 - 2007 Biocceleration Ltd.

OM nucleic - nucleic search, using sw model

Run on: June 14, 2007, 04:31:56; Search time 3358 Seconds

(without alignments)
10931.438 Million cell

updates/sec

Title: US-10-535-378-11

Perfect score: 531 Sequence: 1

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Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 7568541 seqs, 34560148153 residues

Total number of hits satisfying chosen parameters: 15137082

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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2: gb pat:*

3: gb ph:*

4: gb pl:*

5: gb pr:*

6: gb ro:*

7: gb sts:*

98_868.

8: gb_sy:*

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13: gb in:* ·

14: gb om:*

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PRI 22-JUL-1999
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gene, complete
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REFERENCE
          1 (bases 4802 to 9870)
 AUTHORS
          Chang, C.H., Scott, G.K., Kuo, W.L., Xiong, X.,
Suzdaltseva, Y.,
          Park, J.W., Sayre, P., Erny, K., Collins, C., Gray, J.W.
and Benz, C.C.
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ESX: a structurally unique Ets overexpressed early
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   PUBMED
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REFERENCE
            Chang, C.H., Scott, G.K., Baldwin, M.A. and Benz, C.C.
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  TITLE
            Exon 4-encoded acidic domain in the
epithelium-restricted Ets
            factor, ESX, confers potent transactivating capacity
and binds to
            TATA-binding protein (TBP)
            Oncogene 18 (25), 3682-3695 (1999)
  JOURNAL
            10391676
   PUBMED
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REFERENCE
            Chang, C.H., Scott, G.K. and Benz, C.C.
  AUTHORS
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            Direct Submission
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